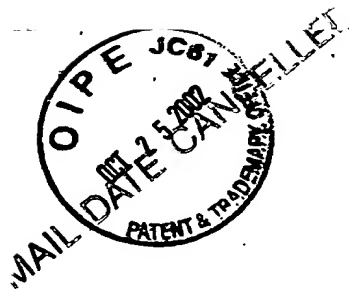


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Figure 1



09881050.112502



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Figure 2

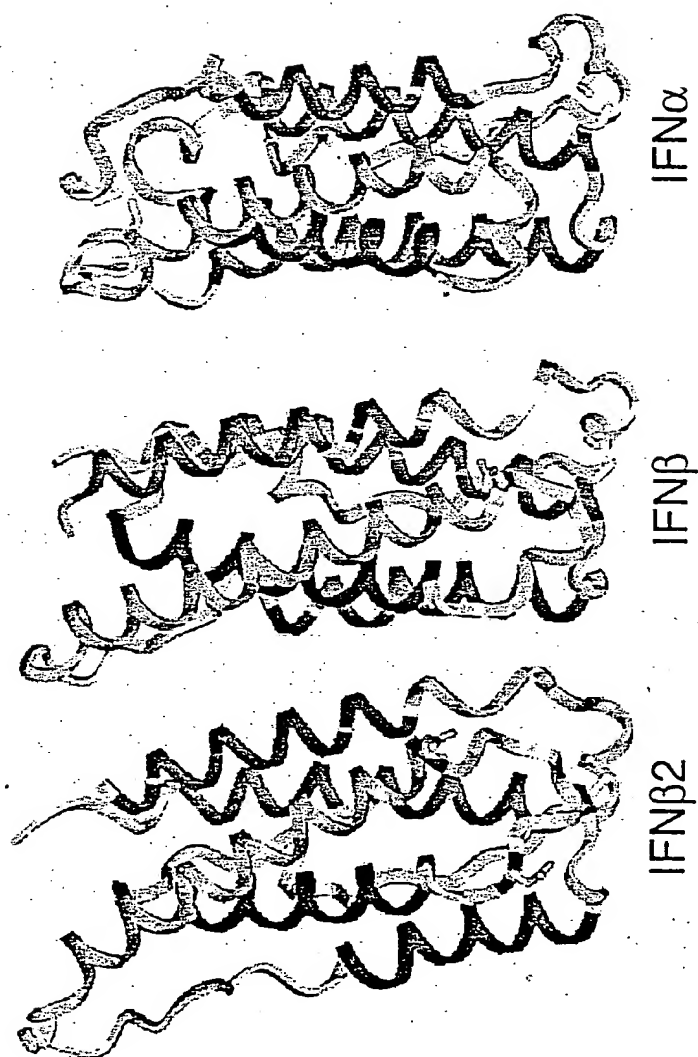


Figure 3



1

IFNB2 (1) MIEKHFFGTVLVLLASTTIFSLDLKIIIFQQRQVNOQSEKILNKIQ-TESIQOCLPHRKNELLPQKSLSP  
 IFNB (1) MTNKCLEQIALLLCFSTALMSYNLLGLQRSSNFQOQKLLWQNGRLEY--CLKDRMNDIPEEIKQL  
 IFNalpha8 (1) MALTFFLLVALVVLVSYSFSSSLGCDLPQTHS-LGNRRATILLAQMR-RISPFSCDKDRHDFEPQEEFDD  
 IFNalpha7 (1) MARSFSLLMVVLVSYSICSLGCDLPQTHS-LPNRRATILLAQMG-RISPFSCDKDRHDFEPQEEFDD  
 IFNalpha6 (1) MALPFALLMALVVLVSCKSSCSLDCDLPQTHS-LGHRRTMMLLAQMR-RISLFSCLDKDRHDFEPQEEFDD  
 IFNalpha5 (1) MALPFVLLMALVVLNCKSSICSLGCDLPQTHS-LSNRRTLMMLAQMG-RISPFSCDKDRHDFEPQEEFDD  
 IFNalpha4b (1) MALSFSLMAVLVLSYSICSLGCDLPQTHS-LGNRRATILLAQMG-RISHFSCDKDRHDFEPQEEFDD  
 IFNalpha21 (1) MALSFSLMAVLVLSYSICSLGCDLPQTHS-LGNRRATILLAQMG-RISPFSCDKDRHDFEPQEEFDD  
 IFNalpha2 (1) MALTEALLVALVLSCKSSCSVGCGLPQTHS-LGSRRTLMMLAQMR-RISLFSCLDKDRHDFEPQEEFDD  
 IFNalpha16 (1) MALSFSLMAVLVLSYSICSLGCDLPQTHS-LGNRRATILLAQMG-RISHFSCDKDRHDFEPQEEFDD  
 IFNalpha14 (1) MALPFALLMALVVLVSCKSSCSLGCNLSQTHS-LNNRRTLMMLAQMR-RISPFSCDKDRHDFEPQEEFDD  
 IFNalpha13 (1) MASPFALLMVLVVLVSCKSSCSLGCGLPETHS-LNNRRTLMMLAQMS-RISPFSSCLMDRHDGFPQEEFDD  
 IFNalpha10 (1) MALSFSLMAVLVLSYSICSLGCDLPQTHS-LGNRRATILLGQMG-RISPFSCDKDRHDFEPQEEFDD  
 IFNomega1 (1) MALFPLLAALVMTSYSVPVSLGCDLPQNHG-LLSRNTLVLLHQM-RISPELCLDKDRHDFEPQEEFDD  
 IFNgamma (1) MKYT-SYTLAFQLCIVLGSGLGCYQDFYVKE----AENLKXYFNAG----H--SDVADNGTLF--LGILK  
 Consensus (1) MAL F LLMALLVLS KS CSLGCDLPQTHS L NRR L LLAQM RISPFSCDKDRHDFEPQEEFDD

70

71

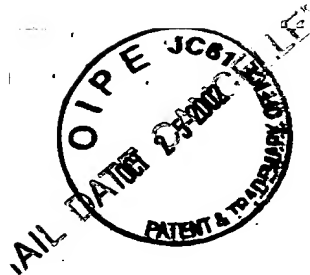
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 IFNB (69) QOQKEDAAALTIYEMLOIFALFRQDSSSTGWNETIVENLLANNVYHQINHLKTVLEEKLEKEDFTRGKLM  
 IFNalpha8 (69) KQFQKAQAISVLHEMIQOTFNLFSTKDSSAALDETLLEDEFYIELDQOLNDLESCVMQEVGVIESPLMYED  
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 IFNalpha21 (69) NQFQKAQAISVLHEMIQOTFNLFSTKDSSATWEQSILKEKFSTELNQOLNDMEACVIOEVGVEETPLMND  
 IFNalpha2 (68) NQFQKAQAISVLHEMIQOTFNLFSTKDSSAAWDETLLDKFYTELYQOLNDLEACVIOEVGVEETPLMND  
 IFNalpha16 (69) NQFQKAQAISVLHEMIQOTFNLFSTKDSSAAWDETLLDKFYTELYQOLNDLEACVIOEVGVEETPLMND  
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 IFNalpha10 (69) NQFQKAQAISVLHEMIQOTFNLFSTEDSSAAWEQSILKEKFSTELYQOLNDLEACVIOEVGVEETPLMND  
 IFNomega1 (69) SOLQKAHVMSVLHEMIQOTFNLFSTEDSSAANMTLLDQHTGLHQOLQHLETCLLQVVGEGESAGAISS  
 IFNgamma (58) NWKEESDRKIMQSCIVSYFYKLFKNFKD---DQS-HQKSVETKEDMN-VKFFNSNKKKRDDFEKITYNY  
 Consensus (71) NQFQKAQAISVLHEMIQOTFNLFSTKDSSAAWDE LDKF TELYQOLNDLEACV QEVGVEETPLMN D

141

IFNB2 (140) LRLOVKMYFRRIHDYLE-NQDYSTCAWATVQVEISRCFLFFVFSLEKLSKQGRPLNDMKQELTTEFRSPR  
 IFNB (139) SSLHLKRYVGRILHYLK-AKEYSHCAWTEVRVEILRNEYFINRLTGYLNRN-----  
 IFNalpha8 (139) SILAVRKYFORITLYLT-EKKYSPCAWEVVRAEIMRSFSLINLQKRLKSKE-----  
 IFNalpha7 (139) FILAVRKYFORITLYLM-EKKYSPCAWEVVRAEIMRSFSLINLQKRLRRKD-----  
 IFNalpha6 (139) SILAVRKYFORITLYLT-EKKYSPCAWEVVRAEIMRSFSSRNQERLRRKE-----  
 IFNalpha5 (139) SILTVRKYFORITLYLT-EKKYSPCAWEVVRAEIMRSFSLSANLQERLRRKE-----  
 IFNalpha4b (139) SILAVRKYFORITLYLT-EKKYSPCAWEVVRAEIMRSFSLINLQKRLRRKD-----  
 IFNalpha21 (139) SILAVRKYFORITLYLT-EKKYSPCAWEVVRAEIMRSFSLKIFQERLRRKE-----  
 IFNalpha2 (138) SILAVRKYFORITLYLK-EKKYSPCAWEVVRAEIMRSFSLSTNLQESLRSKE-----  
 IFNalpha16 (139) SILAVRKYFORITLYLM-GKKYSPCAWEVVRAEIMRSFSLSTNLQKGLRRKD-----  
 IFNalpha14 (139) SILAVRKYFORITLYLM-EKKYSPCAWEVVRAEIMRSFSLSTNLQKRLRRKD-----  
 IFNalpha13 (139) SILAVRKYFORITLYLT-EKKYSPCAWEVVRAEIMRSFSLSTNLQERLRRKE-----  
 IFNalpha10 (139) SILAVRKYFORITLYLI-ERKYSPCAWVVRAEIMRSFSLSTNLQKRLRRKD-----  
 IFNomega1 (139) PALITRRYFQGRIVYLK-EKKYSDCAWEVVRAEIMKSLFLSTNMQERLRSKDRDLGSS-----  
 IFNgamma (122) SVTDENVQRKAHELQVMAELSPAATGKR---KRSQML---FRGRASQ-----  
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210

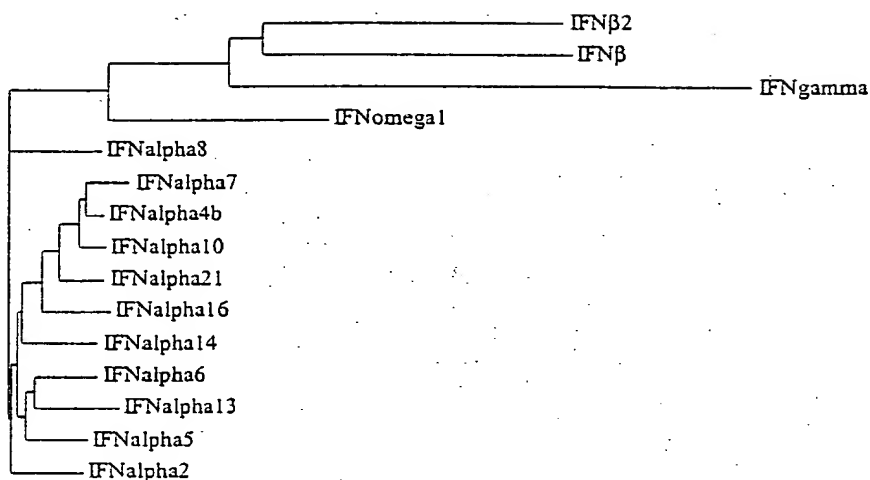
Figure 4



09004050.1125



Protein Level Comparison



Nucleotide Level Comparison

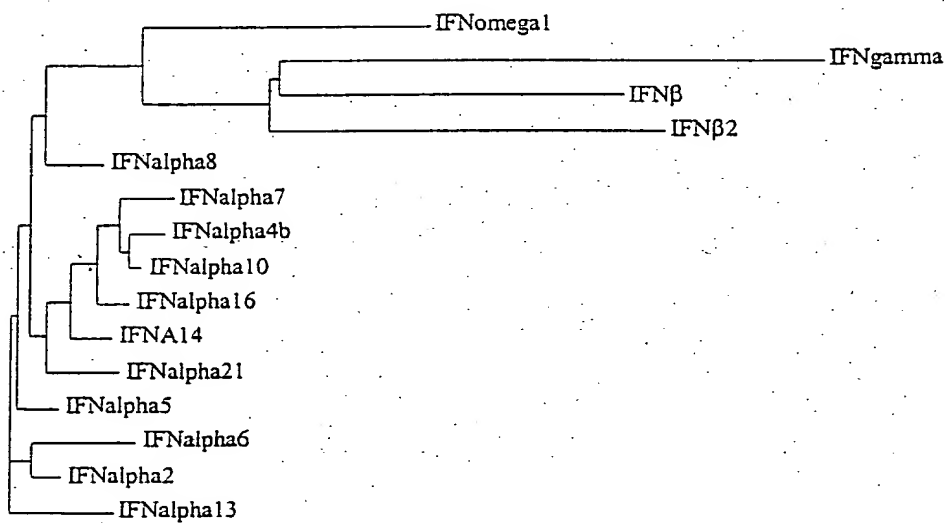


Figure 5

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DATE 25 2002  
PATENT & TRADEMARK

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NOV 25 2002  
TRADEMARK OFFICE

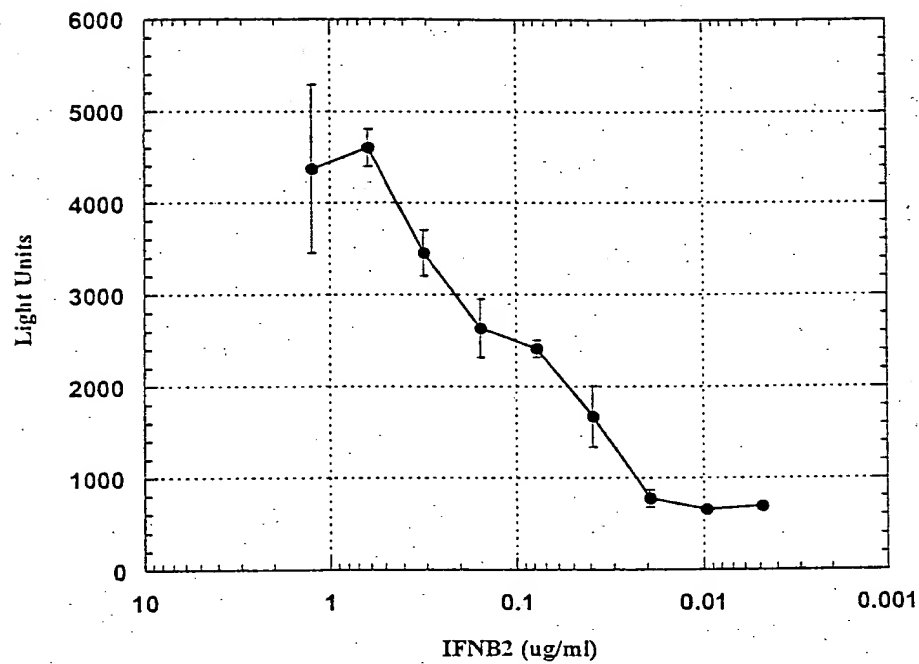


Figure 6



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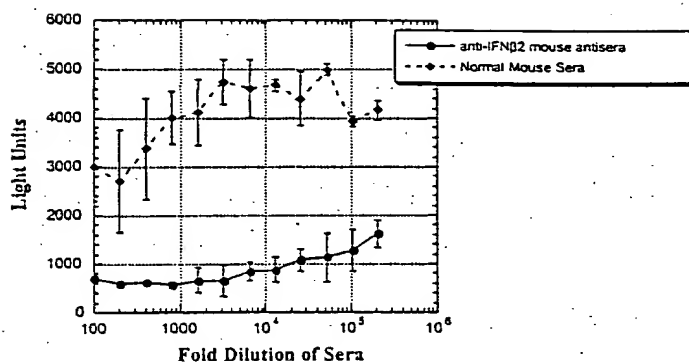
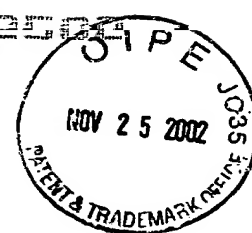


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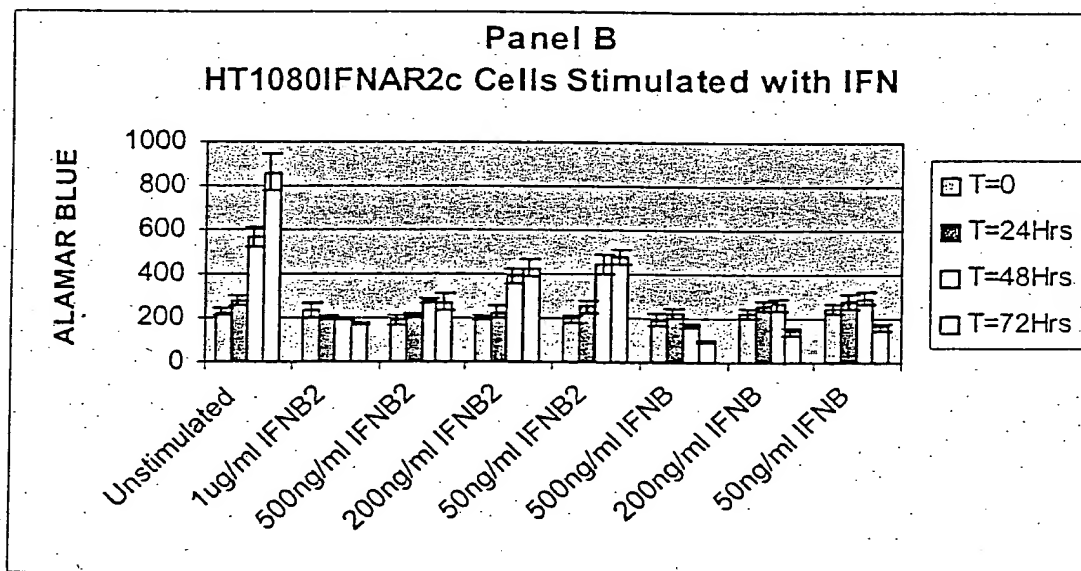
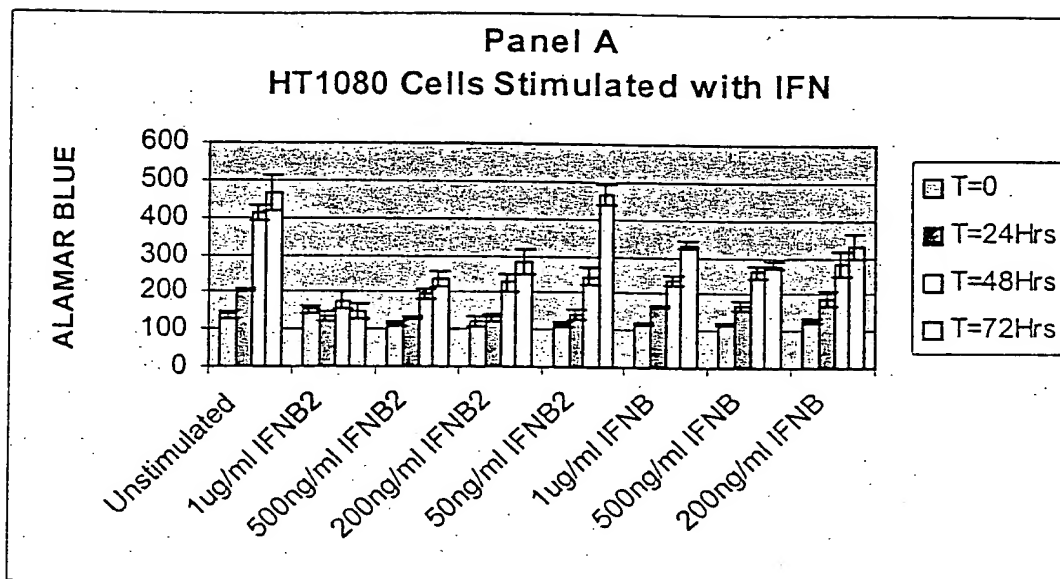
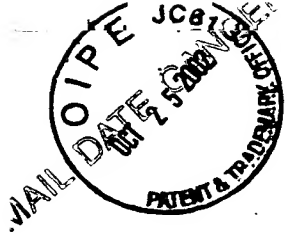


Figure 8





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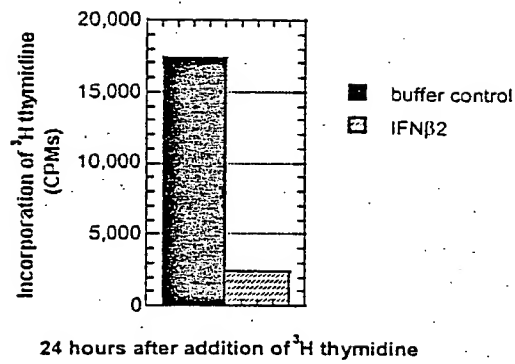
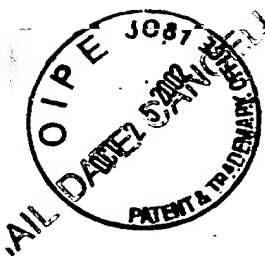


Figure 9



09001050 . 112502

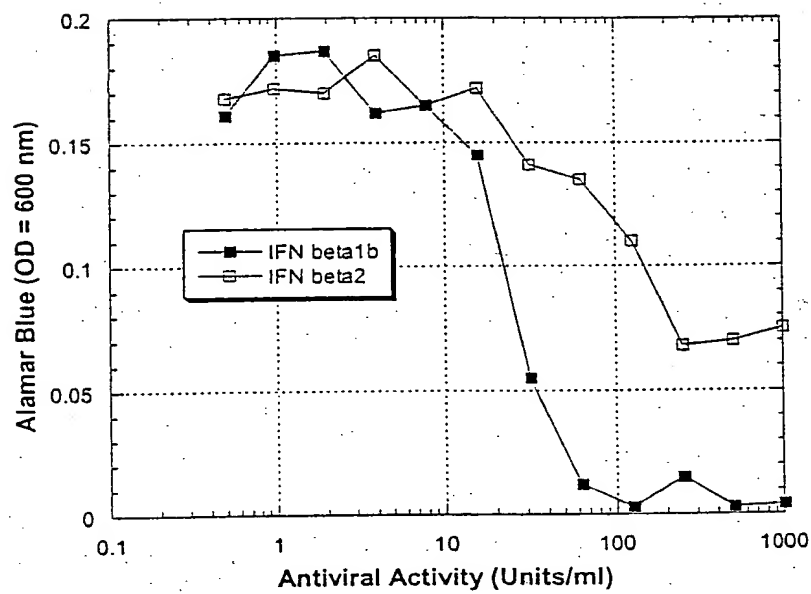
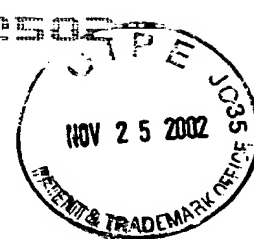


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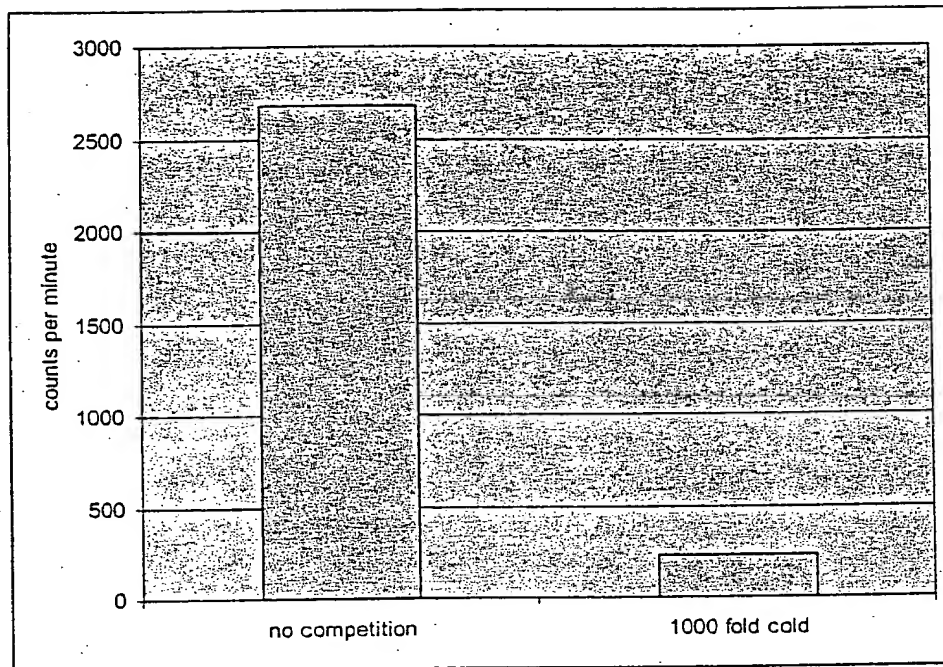
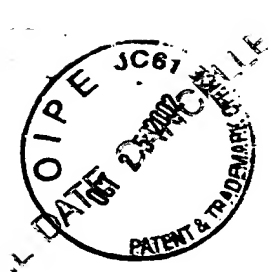
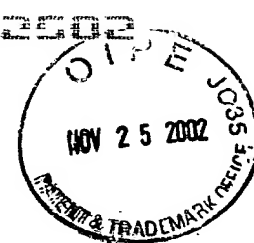
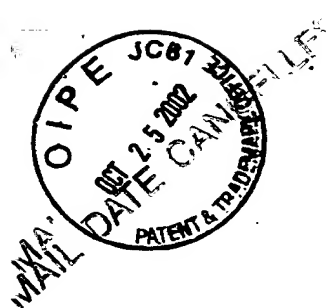


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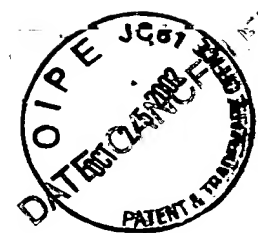
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121   aattgcctca aggaaagcat acaatgaata agttattttg gtacttcctc aaaatagcca  
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241   taatgtgtaa acttgagaa atgacattag aaaacgaaag ctacaaaaga gaacactctt  
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Figure 12



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481 ccagcaaaga caagtgaatc aagaaagttt aaaactcttg aataagttgc aaaccttgtc  
541 aattcagcag tgtctaccac acaggaaaaa cttctctgctt cctcagaagt ctttgagtcc  
601 ttaactgtac caaaaaggac acactctggc cattcttcat gagatgct

Figure 13



09881050 . 112507



MIKHFFGTVLVLLASTTIFSLDLKLIIFQQRQVNQESLKLLNKLQTLISIQ  
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Figure 14

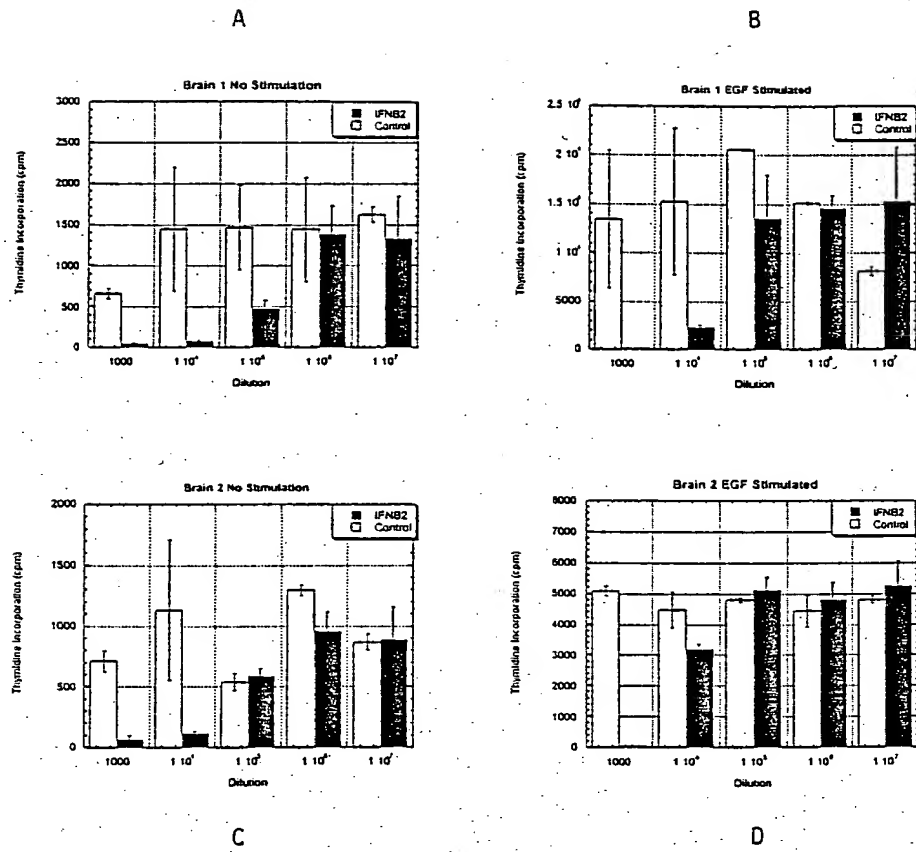
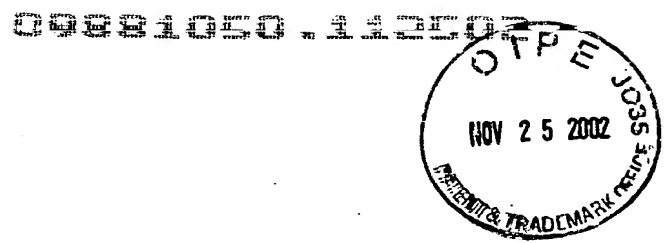
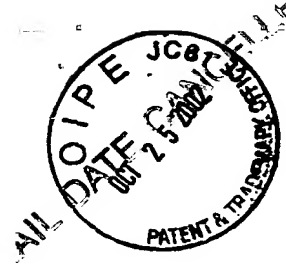


Figure 15